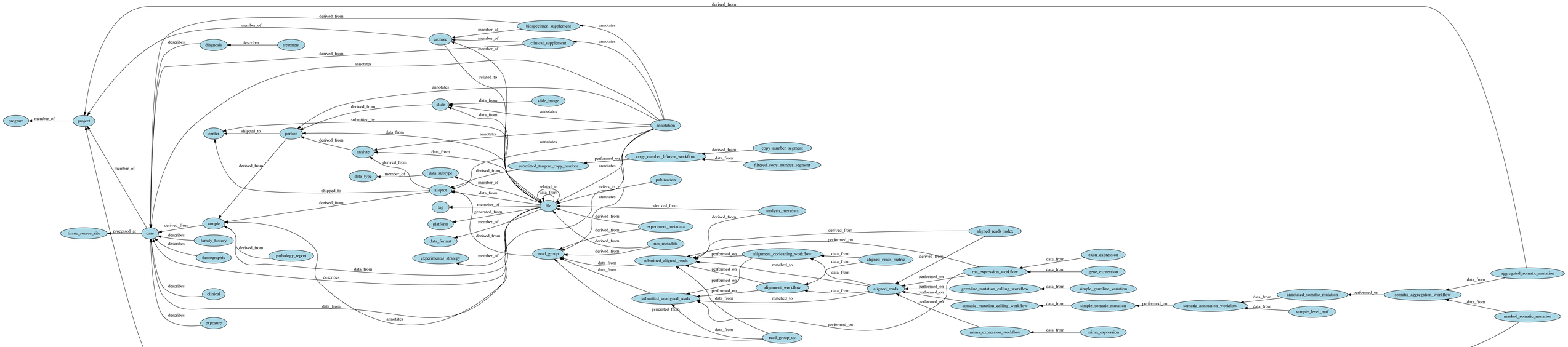
program
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member_of
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biospecimen_supplement
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slide
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annotation
submitted_tangent_copy_number
copy_number_liftover_workflow
copy_number_segment
filtered_copy_number_segment
publication
file
analysis_metadata
experiment_metadata
run_metadata
read_group
submitted_aligned_reads
submitted_unaligned_reads
alignment_cocleaning_workflow
aligned_reads_metric
alignment_workflow
aligned_reads
aligned_reads_index
read_group_qc
rna_expression_workflow
exon_expression
gene_expression
germline_mutation_calling_workflow
simple_germline_variation
somatic_mutation_calling_workflow
simple_somatic_mutation
mirna_expression_workflow
mirna_expression
somatic_annotation_workflow
annotated_somatic_mutation
sample_level_maf
somatic_aggregation_workflow
aggregated_somatic_mutation
masked_somatic_mutation
derived_from
member_of
describes
annotates
related_to
submitted_by
data_from
performed_on
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generated_from
refers_to